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ON protein - protein search, using sw model

Run on: August 28, 2002, 17:31:06 ; Search time 75.04 Seconds

(without alignments)
312.321 Million cell updates/sec

Title: US-09-502-984B-6
Perfect score: 1098
Sequence: 1 KFEKSKALLAARGPPEELCF AEPSEGGFWSAWSEPVSLT 211
Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 747574 seqs, 11073796 residues

Total number of hits satisfying chosen parameters: 747574

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : A_Genesed_032802:*

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 2: /SIDS1/geodata/hold-geneseq/geneseq-emb1/AA1981.DAT:*
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 22: /SIDS1/geodata/hold-geneseq/geneseq-emb1/AA2001.DAT:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result	No.	Score	Query	%	Match Length	DB ID	Description

1	1060	96.5	211	21	AAB21686		Human mature eryth
2	1060	96.5	225	21	AAB21685		Human mature eryth
3	1060	95.5	438	21	AAY41622		Truncated human Epo
4	1060	96.5	488	18	AAW03349		EpoRc fusion prot
5	1060	96.5	503	21	AAB13012		O-tagged erythropo
6	1060	96.5	508	11	ARR0512		EPO receptor. Hom
7	1060	96.5	16	16	ARR70032		Human erythropoietin
8	1060	96.5	508	16	AAR62503		Human erythropoietin
9	1053	95.9	508	16	AAR7518		Human EPO receptor
10	1052	95.8	438	21	RAY44623		R154C truncated hu
11	869.5	79.2	265	15	AAR50326		Mouse soluble EPO

ALIGNMENTS

RESULT 1
ID: AAB21686 standard; peptide: 211 AA.

XX
AC: AAB21686;

XX DT 21-DEC-2000 (first entry)

XX DE Human mature erythropoietin receptor EPOR extracellular domain #2.

XX KW Ligand; cell surface receptor; erythropoietin; EPOR; human;

XX KW protein design automation; PDA.

OS Homo sapiens.

XX PN WO20047612-A2.

PD 17-AUG-2000.

XX PF 11-FEB-2000; 2000WO-US03665.

PR 11-FEB-1999; 99US-0120009.

PR 29-APR-1999; 99US-0131674.

XX PA (XENC-) XENCOR INC.

XX PI Luo, P., Dahlia, B;

XX DR WPI; 2000-549135/50.

PT Screening for ligand analogs and agents which modulate ligand-receptor binding, comprises adding a test ligand to a non-naturally occurring cell surface receptor analog -

XX

EPO receptor seque

MEL EPO receptor.

Mouse erythropoietin

Mouse soluble EPO

Mouse thrombopoietin

Human cell surface

Soluble murine MPL

Type I MPL recepto

Type I MPL r

Mouse type I MPL r

Mouse type I MPL r

Mouse MPL type I r

Murine myeloprolif

Synthetic human er

Synthetic human er

Synthetic human er

Human myeloprolif

Human erythropoietin

Human Zcytov5 vari

PS Example 1; Fig 8; 82pp; English.

XX

CC The present invention relates to a method for screening for a ligand is a mature human erythropoietin receptor (EPOR) extracellular domain.

CC analog, comprising adding a candidate ligand to a non-naturally occurring

CC cell surface receptor analog e.g. erythropoietin receptor (EPOR), and

CC determining the binding of the ligand to the analog. The present sequence

CC is a mature human erythropoietin receptor (EPOR) extracellular domain.

CC Protein Design Automation was carried out on the present sequence, so

CC that it may be used in the present invention as a cell surface receptor

CC analog.

XX

SQ Sequence 211 AA;

Query Match 96.5%; Score 1060; DB 21; Length 211;

Best Local Similarity 93.8%; Pred. No. 2e-106; Matches 198; Conservative 11; Mismatches 2; Indels 0; Gaps 0;

QY 1 KFESKAALLAARGPPEELCFPERLEDVCFEEBEASAGVGGCNFSFOLDEDPWKLCR 60

Db 1 kfeskaallaargpeelcfperledvfcfweeaasagvgghgnysyqledepwklcr 60

OY 61 HOAPRTARGAIREWCSLUTADPSSFVPLRITAASCAPRFVRVHINEVWILDAPVGLA 120

Db 61 hoaprtargatrwcslutaadpssfvplritaaascaprfvrvhinevwildapvglva 120

QY 121 RIADESCHIVWRMLPPETPMTSHIRELDISAGNGAGSVORVELLEGRTCVLSNLGR 180

Db 121 riaedeschivwrmlpppetpmthireldisagngagsvorvellegrtcvlsnlgr 180

QY 121 rlaadesghvvlrwlpppetpmthirelyevdysagngagsvqrveilegtcvlsnlgr 180

Db 121 rlaadesghvvlrwlpppetpmthirelyevdysagngagsvqrveilegtcvlsnlgr 180

QY 181 TRITIAYARMAEPSRGFWSAWSEPVSLT 211

Db 181 tritayarmaepsrgfwswawsepvsilt 211

QY 181 TRITIAYARMAEPSRGFWSAWSEPVSLT 211

Db 190 tritayarmaepsrgfwswawsepvsilt 220

QY 181 TRITIAYARMAEPSRGFWSAWSEPVSLT 211

Db 190 tritayarmaepsrgfwswawsepvsilt 220

RESULT 2

ID AAB21685 standard; peptide; 225 AA.

XX

AC AAB21685;

XX

DT 21-DEC-2000 (first entry)

DE Human mature erythropoietin receptor EPOR extracellular domain #1.

XX

KW Ligand; cell surface receptor; erythropoietin; EPOR; human.

XX

OS Homo sapiens.

XX

PN WO200047612-A2.

XX

PD 17-AUG-2000.

XX

PR 11-FEB-2000; 2000WO-US03665.

XX

PR 11-FEB-1999; 99US-0120009.

PR 29-APR-1999; 99US-0131674.

PA (XENC-) XENCOR INC.

XX

PI Luo P, Dahiyat B;

XX

DR WPI; 2000-549135/50.

PT Screening for ligand analogs and agents which modulate ligand-receptor

PT binding, comprises adding a test ligand to a non-naturally occurring

PT cell surface receptor analog -

XX

Example 1; Fig 8; 82pp; English.

XX

CC The present invention relates to a method for screening for a ligand

CC a cytokine-dependent cell line supports cell population expansion in the

CC absence of exogenous cytokines. Mutant human EPOR is used in treatment of

CC disorders related to inadequate EPOR signalling. The transfected cells

CC may also be used in gene therapy to treat cancer, infectious diseases

CC determining the binding of the ligand to the analog. The present sequence

CC is a mature human erythropoietin receptor (EPOR) extracellular domain.

CC This sequence may be used in the present invention as a cell surface

CC receptor analog.

XX

SQ Sequence 225 AA;

Query Match 96.5%; Score 1060; DB 21; Length 225;

Best Local Similarity 93.8%; Pred. No. 2.2e-106; Matches 198; Conservative 11; Mismatches 2; Indels 0; Gaps 0;

QY 1 KFESKAALLAARGPPEELCFPERLEDVCFEEBEASAGVGGCNFSFOLDEDPWKLCR 60

Db 10 kfeskaallaargpeelcfperledvfcfweeaasagvgghgnysyqledepwklcr 69

QY 61 HOAPRTARGAIREWCSLUTADPSSFVPLRITAASCAPRFVRVHINEVWILDAPVGLA 120

Db 70 hoaprtargatrwcslutaadpssfvplritaaascaprfvrvhinevwildapvglva 129

QY 121 RIADESCHIVWRMLPPETPMTSHIRELDISAGNGAGSVORVELLEGRTCVLSNLGR 180

Db 130 riaedeschivwrmlpppetpmthireldisagngagsvorvellegrtcvlsnlgr 189

QY 181 TRITIAYARMAEPSRGFWSAWSEPVSLT 211

Db 190 tritayarmaepsrgfwswawsepvsilt 220

RESULT 3

ID AAY44622

ID AAY44622 standard; Protein; 438 AA.

XX

AC AAY44622;

XX

DT 07-APR-2000 (first entry)

XX

DE Truncated human EPOR(t439).

XX

KW Truncated human EPOR; erythropoietin receptor; hypersensitive EPOR(t439);

KW mutant human EPOR; EPOR signalling; cancer; infectious disease; HIV;

KW sickle cell anaemia; cytostatic; antimicrobial; antiviral;

KW immunostimulant; anti-anæmic.

XX

OS Homo sapiens.

XX

PN WO9967360-A2.

XX

PD 29-DEC-1999.

XX

PF 25-JUN-1999; 99WO-CA00606.

XX

PR 25-JUN-1998; 98CA-2241576.

PR 25-JAN-1999; 99CA-2260332.

XX

PA (HEMO-) HEMOSOL INC.

XX

PI Bell D, Matthews KE, Mueller SG;

XX

DR WPI; 2000-136979/12.

XX

DR P-PDB; RAZ9634.

XX

PT Serum free defined medium useful for the efficient culture of stem

PT cells used for production of hemoglobin -

XX

PS Example 6; Fig 9; 61pp; English.

XX

CC The present sequence is truncated human EPOR (erythropoietin receptor).

CC Transfection of constitutively active EPOR(t439) by electroporation into

CC a cytokine-dependent cell line supports cell population expansion in the

CC absence of exogenous cytokines. Mutant human EPOR is used in treatment of

CC disorders related to inadequate EPOR signalling. The transfected cells

CC may also be used in gene therapy to treat cancer, infectious diseases

CC (e.g. HIV), sickle cell anemia, and conditions related to abnormal
CC expression of erythropoietin.
XX
SQ Sequence 438 AA;

Query Match 96.5%; Score 1060; DB 21; Length 438;
Best Local Similarity 93.8%; Pred. No. 5.1e-100;
Matches 198; Conservative 11; Mismatches 2; Indels 0; Gaps 0;
Db 34 kfskaalaargpeelcfcftledivcfweeaasagvgpgnysyqlecpwkicrl 93

QY 1 KFESKAALAARGPEELCFCFTLEDIVCFWEAASAGVGPGNYSYQLECPWKICRL 60

QY 61 HOAPTAGRAIRFWCSLPLADTSFVPLRLTAASGARPRFHVIHNEVVLIDAPGVLA 120

Db 94 haptargavrwcslptadtsfvplrltaasagapyrhvihnevvlidapgvla 153

QY 121 RLADESGHIVVIRLPPPTPMHSIRELDISAGNGAGSVQRVELLEGRTCEVLSNRGR 180

Db 154 rlaesghvvlwlppptpmshiryevdvsagnagsvqveilegretcevlsnlgr 213

QY 181 TRITIAYARMARMAPSPFGFWSAWSEPVSLT 211

Db 214 trytfavarmmaepsfsgfwawsepvsilt 244

RESULT 4

ID AAW08349 standard; Protein; 488 AA.

XX AAW08349;

AC

DT 14-MAR-1997 (first entry)

XX

DE EpoRfC fusion protein.

XX

KW Receptor agonist; erythropoietin receptor; EpoR;

KW immunoen; antigen; metallothionein; promoter; IgG1; Fc;

XX anaemia; therapy.

OS Chimeric synthetic.

XX

KEY Location/Qualifiers

FH 1..250

FT /Label= "EpoR-ECD"

FT /note= "erythropoietin receptor extracellular domain"

FT cleavage-site 251..254

FT /note= "Factor Xa cleavage site"

FT Domain 255..488

FT /Label= "Fc domain"

FT /note= "human IgG1 Fc sequence"

PN WO9640231-A1.

XX

PR 19-DEC-1996.

PF 07-JUN-1996; 96WO-US09613.

XX

PR 07-JUN-1995; 95US-0474673.

XX

PA (SMIK) SMITHKLINE BEECHAM CORP.

XX

PI Erickson-Miller CL, Young PR;

XX

DR WPI; 1997-051900/05.

DR N-PSDB; AAT48800.

XX

Recombinant immunogen corresp. to dimeric form of a receptor - used
for generating antibodies able to act as receptor agonists, esp. of
erythropoietin receptor for treating anaemia

XX Example 1; Page 39-41; 83PP; English.

PS

CC A fusion protein (AAW08349) encoded by plasmid mta1sEpoRfC (NAT48800) comprises the human erythropoietin receptor (EpoR) extracellular domain fused (via a Factor Xa cleavage sequence) to the Fc portion of human IgG1. It can be expressed e.g. in transfected Drosophila S2 cells upon induction with copper sulphate. The cells secrete EpoRfC as a dimeric molecule due to the affinity of the Fc moiety for itself. The dimeric receptor can be used as an immunogen to generate antibodies (monoclonal, polyclonal, chimeric, humanised) able to act as EpoR agonists for use in treatment of anaemia.

CC Sequence 488 AA;

Query Match 96.5%; Score 1060; DB 18; Length 488;
Best Local Similarity 93.8%; Pred. No. 5.9e-100;
Matches 198; Conservative 11; Mismatches 2; Indels 0; Gaps 0;
Db 34 kfskaalaargpeelcfcftledivcfweeaasagvgpgnysyqlecpwkicrl 93

QY 61 HOAPTAGRAIRFWCSLPLADTSFVPLRLTAASGARPRFHVIHNEVVLIDAPGVLA 120

Db 94 haptargavrwcslptadtsfvplrltaasagapyrhvihnevvlidapgvla 153

QY 121 RLADESGHIVVIRLPPPTPMHSIRELDISAGNGAGSVQRVELLEGRTCEVLSNRGR 180

Db 154 rlaesghvvlwlppptpmshiryevdvsagnagsvqveilegretcevlsnlgr 213

QY 181 TRITIAYARMARMAPSPFGFWSAWSEPVSLT 211

Db 214 trytfavarmmaepsfsgfwawsepvsilt 244

RESULT 5

ID AAB13012 standard; Protein; 503 AA.

XX

AC AAB13012;

XX

DE 08-DEC-2000 (first entry)

XX

Q-Tagged erythropoietin (EPO) receptor protein.

XX

KW Site specific label; detection; interaction screening; transglutaminase; erythropoietin receptor; EPO.

XX

OS Synthetic.

XX

PN WO200043492-A2.

XX

PD 27-JUL-2000.

XX

PR 20-JAN-2000; 2000WO-US01481.

XX

PR 22-JAN-1999; 99US-0117327.

XX

PA (SMIK) SMITHKLINE BEECHAM CORP.

XX

PT Tew DG, Powell DJ, Meek TD, Chen W;

XX

DR WPI; 2000-49922/44.

XX

PT Screening for a candidate compound for use in bioassays comprises contacting the candidate molecule with a labelled modified protein and detecting the label to identify interaction of the two molecules -

XX

PS Example 4; Page 26; 49PP; English.

XX

CC This invention relates to methods for the site specific modification of

CC a protein, and to a method for screening for a candidate compound which
CC interacts with first protein. The screening method comprises contacting
CC the candidate molecule with a labelled modified first protein and
CC detecting the label to identify interaction of the labelled modified
CC first protein and candidate compound. The first protein is modified to
CC contain a peptide, represented by sequence AAB1305. The method is
CC used to label proteins at specific sites. The present sequence
CC represents a Q-tagged erythropoietin (EPO) receptor constructed in an
CC example of the method of the invention.

XX Sequence 503 AA;

Query Match 96.5%; Score 1060; DB 21; Length 503;

Best Local Similarity 93.8%; Pred. No. 6.1e-106; Matches 198; Conservative 11; Mismatches 2; Indels 0; Gaps 0;

QY 1 KFFSKAALLAARGPEELICFTTERLEDLVCFPPERASAGCAGVPGNFSFSRQLEDEPWKLCRL 60

Db 34 kfeskallaargpeelicftterleldivcfweeaasqygggnfsfsrqledepwkicrl 93

QY 61 HQAPTARGAIRFWCSLPLADTSFVPLRLTAASGAPRFRVTHNEVVLIDAPVLA 120

Db 94 hqaptargavrwcslpladtsfvplrltaasgapyrhvihnevildapvla 153

QY 121 RLADESHWVIRMLPPETPMWSHIRELDISAGNGAGSVQRVELGRTECVLSNRGR 180

Db 154 rladesgqvirwipppetpmthiryevdvsagnagsvqrveilegrtecvlsnlg 213

QY 181 TRITIAYARMAPSPSGFGFWSAWSEPSLLT 211

Db 214 trytfavrmarepsfgfwawsawsepsllt 244

XX Sequence 508 AA;

Query Match 96.5%; Score 1060; DB 11; Length 508;

Best Local Similarity 93.8%; Pred. No. 6.2e-106; Matches 198; Conservative 11; Mismatches 2; Indels 0; Gaps 0;

QY 1 KFFSKAALLAARGPEELICFTTERLEDLVCFPPERASAGCAGVPGNFSFSRQLEDEPWKLCRL 60

Db 34 kfeskallaargpeelicftterleldivcfweeaasqygggnfsfsrqledepwkicrl 93

QY 61 HQAPTARGAIRFWCSLPLADTSFVPLRLTAASGAPRFRVTHNEVVLIDAPVLA 120

Db 94 hqaptargavrwcslpladtsfvplrltaasgapyrhvihnevildapvla 153

QY 121 RLADESHWVIRMLPPETPMWSHIRELDISAGNGAGSVQRVELGRTECVLSNRGR 180

Db 154 rladesgqvirwipppetpmthiryevdvsagnagsvqrveilegrtecvlsnlg 213

QY 181 TRITIAYARMAPSPSGFGFWSAWSEPSLLT 211

Db 214 trytfavrmarepsfgfwawsawsepsllt 244

RESULT 6

AAR06512

ID AAR06512 standard; protein; 508 AA.

XX

AC AAR06512;

XX

DT 04-JAN-1991 (first entry)

XX

DE EPO receptor.

DE

XX

KW Erythropoietin; Diamond Blackfan anaemia; polycythemia vera.

XX

OS Homo sapiens.

XX

PN WO9008842-A.

XX

PD 09-AUG-1990.

XX

PF 01-FEB-1990; 90WO-US00635.

XX

PR 03-FEB-1989; 89US-0306503.

XX

PA (GENE-) GENETICS INST INC.

PA (WHIT-) WHITEHEAD INST.

XX

PI Dr'andrea A., Wong G;

XX

DR WPI: 1990-260931/34.

XX

DR N-PSDB; AAQ05748.

PT Erythropoietin receptor and gene - used for developing reagents

PT and systems to control and study erythropoiesis.

XX

PS Disclosure; Fig 2; 53pp; English.

XX

CC The sequence was deduced from DNA obtd. from a clone isolated from

CC a commercially available human genomic cDNA library in Phage

CC Lambda Fix (Stratagene). The sequence encodes a type I trans-

membrane protein with binding affinity for EPO. The gene and

CC recombinant EPO receptor produced on expression of the DNA are
CC used to develop reagents and systems to control and study
CC erythropoiesis. It is believed that the EPO receptor is dys-
CC functional in individuals with Diamond Blackfan anaemia, and may
CC be hyperactive in polycythemia vera.
CC See also AAR06511 (murine EPO receptor).

RESULT 7

AAR70032

ID AAR70032 standard; Protein; 508 AA..

XX

AC AAR70032;

XX

DT 07-OCT-1995 (first entry)

XX

DE Human erythropoietin receptor.

XX

KW Erythropoietin receptor; extracellular domain.

XX

OS Homo sapiens.

XX

FH Key Location/Qualifiers

FT Domain /note= "extracellular domain"

FT Domain 9..83

FT Misc-difference /note= "extracellular domain"

FT Misc-difference 25..29

FT Misc-difference /note= "forward primer AAQ82991 specific site"

FT Misc-difference 222..226

FT /note= "reverse primer AAQ82992 specific site"

XX

PN WO9505469-A.

XX

PD 23-FEB-1995.

XX

PF 15-AUG-1994; 94WO-US09298.

XX

PR 16-AUG-1993; 93US-0106815.

XX

PA (IEEJ/) LEE J Y.

XX

PI Lee JY;

XX

DR WPI: 1995-098767/13.

XX

DR N-PSDB; AAQ02990.

XX

New pure human erythropoietin receptor fragment - obtd. by

PT expression as a fusion protein having a thrombin proteolytic

PT cleavage site.

PA	(CHIL-) CHILDRENS MEDICAL CENT.	XX	PN	MD9957360-A2.
(GENE) GENETICS INST INC.		XX	PD	29-DEC-1999.
(WHED) WHITEHEAD INST BIOMEDICAL RES.		XX	PF	25-JUN-1999; 99WO-CA00606.
D'andrea A, Jones SS, Wong GG;		XX	PR	25-JUN-1998; 98CA-2241576.
DR	WPI: 1994-025409/03	XX	PR	25-JAN-1999; 99CA-2260332.
XX	N-FSDB; AAQ5395.	PT	PI	D'andrea A, Jones SS, Wong GG;
PS	Recombinant DNA encoding erythropoietin receptor - used to develop probe for study, treatment or diagnosis of disorders in which receptor is dysfunctional	XX	PT	WPI: 1994-025409/03
XX	Disclosure: Fig 9; 24pp; English.	XX	PT	N-FSDB; AAQ5395.
CC	Mouse erythroleukaemia (MEL) cells were used to construct a cDNA library. The cDNA was used to transfect COS-1 cells and these were screened for radiiodinated erythropoietin (EPO) binding to isolate cDNA encoding the EPO receptor. This cDNA was used as a probe to screen a human genomic cDNA library to obtain DNA encoding the human EPO receptor. The cDNA may be used to study, treat or diagnose disorders in which the EPO receptor is dysfunctional. The EPO receptor may also be used to raise antibodies for treating hypersensitivity to EPO or who have elevated levels of EPO. The word is pref. used for treating anaemias, primary proliferative polycythaemia and secondary polycythaemia. See also AAR7517.	CC	PS	WPI: 2000-136979/12.
CC	Sequence 508 AA;	XX	DR	N-PSDB; AAZ49636.
Query Match 95.9%; Score 1053; DB 15; Length 508; Best Local Similarity 92.9%; Pred. No. 3.6e-105; Matches 196; Conservative 12; Mismatches 3; Indels 0; Gaps 0;	XX	XX	XX	Example 6; Fig 10; 61pp; English.
OY 1 KFESKAALIAARSPPEELCFTTERELDVCFEEAASAGVGPGNFSFSQLEDEPWKICRL 60 34 kfeskaalaaargpeelcfttereldvicfwegasagvgpgnfsqledepwkicrl 93	XX	PT	CC	The present sequence is R154C truncated human EpoR (erythropoietin receptor). Transfection of constitutively active EpoR(t439; R154C) by electroporation into a cytokine-dependent cell line supports cell population expansion in the absence of exogenous cytokines. Mutant human EpoR is used in treatment of disorders related to inadequate EpoR signalling. The transfected cells may also be used in gene therapy to treat cancer, infectious diseases (e.g. HIV), sickle cell anaemia, and conditions related to abnormal expression of erythropoietin.
Db 61 HQAPTARGAIRFWCSLPPADTSSFPVPLRLTAASGAPRFHRVTHINEWVLIDAPVGLVA 120 94 hqptargavrfwcslptadtsfpvplrltaasgapyrhvihnevildapvgiva 153	XX	PS	CC	Sequence 438 AA;
OY 61 HQAPTARGAIRFWCSLPPADTSSFPVPLRLTAASGAPRFHRVTHINEWVLIDAPVGLVA 120 34 kfeskaalaaargpeelcfttereldvicfwegasagvgpgnfsqledepwkicrl 93	XX	XX	CC	Query Match 95.8%; Score 1052; DB 21; Length 438; Best Local Similarity 93.4%; Pred. No. 3.8e-105; Matches 197; Conservative 11; Mismatches 3; Indels 0; Gaps 0;
Db 94 hqptargavrwwcslptadtsfpvplrltaasgapyrhvihnevildapvgiva 153 121 RLADESGHVVIRWLPPPETPMWSHIRFIELDISAGNGAGSVQRVELLEBRCTECVLSNLRGR 180	XX	PT	CC	Query Match 95.8%; Score 1052; DB 21; Length 438; Best Local Similarity 93.4%; Pred. No. 3.8e-105; Matches 197; Conservative 11; Mismatches 3; Indels 0; Gaps 0;
Db 154 rladesgnvvirwlpppetpmshiryevdvsagngagsvqrveilegrtecvlsnlggr 213 OY 181 TRITIATVRMARPSPSGFWSANSEPVSLT 211	XX	PS	CC	Query Match 95.8%; Score 1052; DB 21; Length 438; Best Local Similarity 93.4%; Pred. No. 3.8e-105; Matches 197; Conservative 11; Mismatches 3; Indels 0; Gaps 0;
Db 214 trytfavrmaraepsfggfwawsawsepvsllt 244 RESULT 10	XX	XX	CC	Query Match 95.8%; Score 1052; DB 21; Length 438; Best Local Similarity 93.4%; Pred. No. 3.8e-105; Matches 197; Conservative 11; Mismatches 3; Indels 0; Gaps 0;
AAV44623 standard; Protein: 438 AA.	XX	XX	AC	Query Match 95.8%; Score 1052; DB 21; Length 438; Best Local Similarity 93.4%; Pred. No. 3.8e-105; Matches 197; Conservative 11; Mismatches 3; Indels 0; Gaps 0;
AC AAV44623;	XX	XX	AC	Query Match 95.8%; Score 1052; DB 21; Length 438; Best Local Similarity 93.4%; Pred. No. 3.8e-105; Matches 197; Conservative 11; Mismatches 3; Indels 0; Gaps 0;
DT 07-APR-2000 (first entry)	XX	XX	AC	Query Match 95.8%; Score 1052; DB 21; Length 438; Best Local Similarity 93.4%; Pred. No. 3.8e-105; Matches 197; Conservative 11; Mismatches 3; Indels 0; Gaps 0;
XX R154C truncated human EpoR(t439).	XX	XX	AC	Query Match 95.8%; Score 1052; DB 21; Length 438; Best Local Similarity 93.4%; Pred. No. 3.8e-105; Matches 197; Conservative 11; Mismatches 3; Indels 0; Gaps 0;
DE R154C truncated human EpoR(t439).	XX	XX	AC	Query Match 95.8%; Score 1052; DB 21; Length 438; Best Local Similarity 93.4%; Pred. No. 3.8e-105; Matches 197; Conservative 11; Mismatches 3; Indels 0; Gaps 0;
XX Truncated human EpoR; erythropoietin receptor; hypersensitive EpoR(t439); mutant human EpoR; EpoR signalling; cancer; infectious disease; HIV; sickle cell anaemia; cytostatic; antimicrobial; antiviral; immunomodulant; anti-anemic.	XX	XX	AC	Query Match 95.8%; Score 1052; DB 21; Length 438; Best Local Similarity 93.4%; Pred. No. 3.8e-105; Matches 197; Conservative 11; Mismatches 3; Indels 0; Gaps 0;
OS Homo sapiens.	OS	OS	AC	Query Match 95.8%; Score 1052; DB 21; Length 438; Best Local Similarity 93.4%; Pred. No. 3.8e-105; Matches 197; Conservative 11; Mismatches 3; Indels 0; Gaps 0;
FH Key difference 154 /note= "Wild type Arg substituted by Cys"	FH	FH	AC	Query Match 95.8%; Score 1052; DB 21; Length 438; Best Local Similarity 93.4%; Pred. No. 3.8e-105; Matches 197; Conservative 11; Mismatches 3; Indels 0; Gaps 0;
FT Misc-difference 154 /note= "Wild type Arg substituted by Cys"	FT	FT	AC	Query Match 95.8%; Score 1052; DB 21; Length 438; Best Local Similarity 93.4%; Pred. No. 3.8e-105; Matches 197; Conservative 11; Mismatches 3; Indels 0; Gaps 0;

FT Protein	26..205	FT Domain	272..507
XX /note= "Mature EPO-R fragment"		FT /label=intracellular domain	
PN JP06038787-A.		FT Modified-site	75..77
XX PD 15-FEB-1994.		FT Modified-site	/label=N-linked_glycos
XX PF 04-MAR-1992; 92JP-0082865.		FT /label=N-linked_glycos	182..184
XX PR 04-MAR-1992; 92JP-0082865.		FT /label=N-linked_glycos	
XX PA (SNOW) SNOW BRAND MILK PROD CO LTD.		PR 01-FEB-1990; 90WO-US00635.	
XX DR WPI; 1994-034847/12.		PR 03-FEB-1989; 89US-0306503.	
XX N-PSDB; AAQ04853.		PA (GENE-) GENETICS INST INC.	
XX Soluble erythropoietin receptor protein - and DNA coding for		PA (WHIT-) WHITEHEAD INST.	
PT SEPO-R, useful as diagnostic reagent		PT D'andrea A, Wong G;	
XX disclosure; Page 5-6; 9pp; Japanese.		DR WPI; 1990-260931/34.	
CC This sequence represents a fragment of the murine soluble erythro-		DR N-PSDB; AAQ05747.	
CC poietin (EPO) receptor protein (SEPO-R). This protein is able to		XX PT Erythropoietin receptor and gene - used for developing reagents	
CC bind to EPO and has antigenic properties. The molecular		CC and systems to control and study erythropoiesis.	
CC weight of the full length protein is pref 33 or 29 kD. The protein		CC See also AAR06512 (human EPO receptor).	
CC is useful as a drug, as a diagnostic agent and a biochemical reagent.		XX PS Disclosure; Fig 1; 53pp; English.	
SQ Sequence 265 AA;		XX CC The sequence was deduced from DNA from a clone isolated from a	
Query Match 79.2%; Score 859.5; DB 15; Length 265;		CC cDNA library prep. from uninduced murine erythroleukemia cells.	
Best Local Similarity 77.7%; Pred. No. 1.1e-85;		CC It is a type I transmembrane protein with binding affinity for EPO.	
Matches 164; Conservative 22; Mismatches 24; Indels 1; Gaps 1;		CC The gene and recombinant EPO receptor produced on expression of	
QY 1 KFESKAALIARAOPPEELCFTERLELDVCFERRASAGNGAGSNSFQLEDEPWKLICRL 60		CC the DNA are used to develop reagents and systems to control and	
Db 34 kfeskaalilasqsellcftrledlvclfweeaassgm-dtnysfyqleyesrkcs1 92		CC study erythropoiesis. It is believed that the EPO receptor is	
QY 61 HQAPTAARGAIRFWCSLPLADTSSFPVPLERLRTASGAPRFRHVTIHNEVLLDAPVGLA 120.		CC dysfunctional in individuals with Diamond Blackfan anaemia, and	
Db 93 hqaptvqsvrfwclspladtssfpvlelqvteasgspryhrnihnevildapglia 152		CC may be hyperactive in polycythaemia vera.	
Query Match 79.2%; Score 859.5; DB 11; Length 507;		XX SQ Sequence 507 AA;	
Best Local Similarity 77.7%; Pred. No. 2.5e-85;		CC The sequence was deduced from DNA from a clone isolated from a	
Matches 164; Conservative 22; Mismatches 24; Indels 1; Gaps 1;		CC cDNA library prep. from uninduced murine erythroleukemia cells.	
Db 153 rraegshvvlrpplggapmtthirevdvsagnragtqrvvelegrtcvlsnlgg 212		CC It is a type I transmembrane protein with binding affinity for EPO.	
QY 181 TRTIAVARMARPSFEGFWSAWSEPVSLT 211		CC The gene and recombinant EPO receptor produced on expression of	
Db 213 trytfavarmepsfsfqfwawsepasl1 243		CC the DNA are used to develop reagents and systems to control and	
RESULT 12		CC study erythropoiesis. It is believed that the EPO receptor is	
AAR06511 ID AAR06511 standard; protein; 507 AA.		CC dysfunctional in individuals with Diamond Blackfan anaemia, and	
XX AC		CC may be hyperactive in polycythaemia vera.	
XX AAR06511;		XX SQ Sequence 507 AA;	
XX 04-JAN-1991 (first entry)		CC The sequence was deduced from DNA from a clone isolated from a	
XX DE EPO receptor sequence deduced from DNA of clone 190.		CC cDNA library prep. from uninduced murine erythroleukemia cells.	
XX KW Erythropoietin; diamond Blackfan anaemia; polycythemia vera.		CC It is a type I transmembrane protein with binding affinity for EPO.	
OS Mus musculus.		CC The gene and recombinant EPO receptor produced on expression of	
XX Key location/Qualifiers		CC the DNA are used to develop reagents and systems to control and	
FT Peptide 1..24		CC study erythropoiesis. It is believed that the EPO receptor is	
FT /label=signal peptide 25..248		CC dysfunctional in individuals with Diamond Blackfan anaemia, and	
FT /label=extracellular domain /note=EPO binding region 248..271		CC may be hyperactive in polycythaemia vera.	
FT Domain /label=transmembrane domain		XX RESULT 13	
FT AAR47517 standard; Protein; 507 AA.		XX AAR47517	
XX AC AAR47517;		XX DT 24-JUN-1994 (first entry)	
XX DE MEL EPO receptor.		XX DE MEL EPO receptor.	
KW Erythropoietin receptor; recombinant; murine; anaemia.			

XX	OS	Mus musculus.	RESULT 14
FH	Key	Location/Qualifiers	AAR69502
FT	Peptide	1..24	ID AAR69502 standard; Protein: 507 AA.
FT	Protein	/note= "signal"	XX
FT	Modified-site	25..507	AC AAR69502;
FT	Modified-site	75	XX
FT	Region	/note= "mature EPO receptor"	DT 10-AUG-1995 (first entry)
FT	Region	383	DE Mouse erythropoietin receptor.
FT	Region	250..271	XX
FT	Region	/note= "potential N-glycosylation site"	KW Erythropoietin receptor; anemia therapy; signal peptide;
PN	US5278065-A.	XX	KW transmembrane region; N-linked glycosylation.
XX	OS	Mus musculus.	OS
XX	PD	11-JAN-1994.	XX
XX	FH	Key	Key
PF	Peptide	1..24	Location/Qualifiers
XX	PF	/note= "signal peptide"	FT
PR	PR	03-FEB-1989; 89US-0306503.	FT Protein
PR	PR	25-MAR-1991; 91US-0678877.	FT /note= "mature protein"
XX	PA	(CHIL-) CHILDRENS MEDICAL CENT.	FT Modified-site
PA	PA	{GEMY } GENETICS INST INC.	FT 75..77 /note= "N-linked glycosylation site"
PA	PA	{WHED } WHITEHEAD INST BIOMEDICAL RES.	FT Domain 250..271 /note= "transmembrane region"
XX	PI	D'andrea A, Jones SS, Wong GG;	FT Modified-site 383..385 /note= "N-linked glycosylation site"
XX	DR	WPI: 1994-025409/03.	FT
XX	DR	N-PSDB; AAQ23994.	PN US5378808-A.
XX	XX	XX	XX
PT	PT	Recombinant DNA encoding erythropoietin receptor - used to develop prods. for study, treatment or diagnosis of disorders in which receptor is dysfunctional.	PD 03-JAN-1995.
PT	PS	Disclosure; Fig 2; 24pp; English.	XX
PT	XX	Mouse erythroleukaemia (MEL) cells were used to construct a cDNA library. The cDNA was used to transfect COS-1 cells and these were screened for radiolabelled erythropoietin (EPO) binding to isolate cDNA encoding the EPO receptor. The cDNA may be used to isolate the EPO receptor from other sources and to study, treat or diagnose disorders in which the EPO receptor is dysfunctional. The EPO receptor may also be used to raise antibodies or for treating hyporesponsivity to EPO or who have elevated levels of EPO. The prod. is pref. used for treating anemias, primary proliferative polycythemia and secondary polycythemia. See also AAR47518.	PF 03-FEB-1989; 89US-0306503.
CC	CC	CC	PR 03-FEB-1989; 89US-0306503.
CC	CC	CC	PR 25-MAR-1991; 91US-0678877.
CC	CC	CC	PR 10-JUN-1993; 93US-0075069.
CC	XX	XX	XX
CC	PA	(GEMY) GENETICS INST INC.	PA
CC	PI	D'andrea A, Jones SS, Wong GG;	PA
CC	DR	WPI: 1995-051310/07.	PA
CC	DR	N-PSDB; AAQ18191.	XX
CC	PT	New recombinant erythropoietin receptor polypeptide(s) - used for detection, purification, and therapy and for prodn. of antibodies for detection and therapy	PT
CC	XX	XX	PT
PS	PS	Claim 1; Fig 2; 24pp; English.	XX
XX	CC	The sequence corresponds to a mouse erythropoietin receptor, including putative signal peptide and transmembrane regions, and 2 N-linked glycosylation sites. The protein is derived from mouse erythroleukemia cells and may be used in drug design or in pharmaceutical compositions for therapy of anemia.	CC
SQ	CC	Sequence 507 AA;	CC
SQ	XX	The sequence corresponds to a mouse erythropoietin receptor, including putative signal peptide and transmembrane regions, and 2 N-linked glycosylation sites. The protein is derived from mouse erythroleukemia cells and may be used in drug design or in pharmaceutical compositions for therapy of anemia.	XX
SQ	XX	Sequence 507 AA;	XX
Query Match	CC	79.2%; Score 869.5; DB 15; Length 507;	Query Match
Best Local Similarity	CC	77.7%; Pred. No. 2.5e-85;	Best Local Similarity
Matches	CC	Mismatches 22; Indels 1; Gaps 1;	Matches 164; Conservative 22; Mismatches 24; Indels 1; Gaps 1;
QY	1	KFESKAALLAARGPPEELICFTTERLEDIVCFEREEASAGVGPGNFNSFSFOLEDEPWKLCR 60	QY 1 KFESKAALLAARGPPEELICFTTERLEDIVCFEREEASAGVGPGNFNSFSFOLEDEPWKLCR 60
Db	34	kfeskaallassgseelclftqiledivcfweeaasggm-dfnstsyqlegeskrscl 92	Db 34 kfeskaallassgseelclftqiledivcfweeaasggm-dfnstsyqlegeskrscl 92
Db	61	HQAPTAARGAIRFWCSLPRADSSFPVPLRLIAASGAPRFHRVHNEVWLIDAPVGLVA 120	Db 34 kfeskaallassgseelclftqiledivcfweeaasggm-dfnstsyqlegeskrscl 92
QY	61	HQAPTAARGAIRFWCSLPRADSSFPVPLRLIAASGAPRFHRVHNEVWLIDAPVGLVA 120	QY 1 KFESKAALLAARGPPEELICFTTERLEDIVCFEREEASAGVGPGNFNSFSFOLEDEPWKLCR 60
Db	93	hqaptvrgsvrwsplbtadtsffvplqteaspspryhrnihnevvlidapaglia 152	Db 34 kfeskaallassgseelclftqiledivcfweeaasggm-dfnstsyqlegeskrscl 92
QY	121	RKLDERSHWVWLPPEPPTMPSHIREELDSAGANGAGSVQWELLEGRTCEQLWSNLRG 180	QY 1 KFESKAALLAARGPPEELICFTTERLEDIVCFEREEASAGVGPGNFNSFSFOLEDEPWKLCR 60
Db	153	rraeegshvrlwlpqpgapmtthiryevdvsagrarrgtqrrevilegtcvlsnrg 212	Db 34 kfeskaallassgseelclftqiledivcfweeaasggm-dfnstsyqlegeskrscl 92
QY	181	TRTIAYVARMABPSSEGFWSAWSEPVSLT 211	QY 61 HQAPTAARGAIRFWCSLPTADSSFPVPLRLIAASGAPRFHRVHNEVWLIDAPVGLVA 120
Db	213	tlytfavarnmaepsifgrwsawsepaslt 243	Db 34 kfeskaallassgseelclftqiledivcfweeaasggm-dfnstsyqlegeskrscl 92

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